

RAW SEQUENCE LISTING  
PATENT APPLICATION US/08/951,733

DATE: 05/21/98  
TIME: 12:38:31

INPUT SET: S25958.raw

This Raw Listing contains the General  
Information Section and up to the first 5 pages.

SEQUENCE LISTING

ENTERED

- 1
- 2
- 3 (1) General Information:
- 4
- 5 (i) APPLICANT: Harrington, Lea A.
- 6 Robinson, Murray O.
- 7
- 8 (ii) TITLE OF INVENTION: NOVEL GENES ENCODING TELOMERASE PROTEINS
- 9
- 10 (iii) NUMBER OF SEQUENCES: 44
- 11
- 12 (iv) CORRESPONDENCE ADDRESS:
- 13 (A) ADDRESSEE: Amgen Inc.
- 14 (B) STREET: One Amgen Center Drive
- 15 (C) CITY: Thousand Oaks
- 16 (D) STATE: CA
- 17 (E) COUNTRY: USA
- 18 (F) ZIP: 91320-1789
- 19
- 20 (v) COMPUTER READABLE FORM:
- 21 (A) MEDIUM TYPE: Floppy disk
- 22 (B) COMPUTER: IBM PC compatible
- 23 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
- 24 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- 25
- 26 (vi) CURRENT APPLICATION DATA:
- 27 (A) APPLICATION NUMBER: US 08/951,733
- 28 (B) FILING DATE: 16-OCT-1997
- 29 (C) CLASSIFICATION:
- 30
- 31 (vii) PRIOR APPLICATION DATA:
- 32 (A) APPLICATION NUMBER: US 08/873,039
- 33 (B) FILING DATE: 11-JUN-1997
- 34
- 35 (vii) PRIOR APPLICATION DATA:
- 36 (A) APPLICATION NUMBER: US 08/751,189
- 37 (B) FILING DATE: 15-NOV-1996
- 38
- 39 (viii) ATTORNEY/AGENT INFORMATION:
- 40 (A) NAME: Oleski, Nancy A.
- 41 (B) REGISTRATION NUMBER: 34,688
- 42 (C) REFERENCE/DOCKET NUMBER: A-433B
- 43
- 44 (ix) TELECOMMUNICATION INFORMATION:
- 45 (A) TELEPHONE: (805) 447-6504
- 46 (B) TELEFAX: (805) 499-8011

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## (2) INFORMATION FOR SEQ ID NO:1:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 7881 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

## (ii) MOLECULE TYPE: cDNA

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

ATGGAAAAAC TCCATGGGCA TGTGTCTGCC CATCCAGACA TCCTCTCCTT GGAGAACCGG	60
TGCCTGGCTA TGCTCCCTGA CTTACAGCCC TTGGAGAAAC TACATCAGCA TGTATCTACC	120
CACTCAGATA TCCTCTCCTT GAAGAACCAG TGCCTAGCCA CGCTTCCTGA CCTGAAGACC	180
ATGGAAAAAC CACATGGATA TGTGTCTGCC CACCCAGACA TCCTCTCCTT GGAGAACCAG	240
TGCCTGGCCA CACTTCTCTGA CCTGAAGACC ATGGAGAAAC CACATGGACA TGTCTCTGCC	300
CACCCAGACA TCCTCTCCTT GGAGAACCGG TGCCTGGCCA CCCTCCCTAG TCTAAAGAGC	360
ACTGTGTCTG CCAGCCCCCTT GTTCCAGAGT CTACAGATAT CTCACATGAC GCAAGCTGAT	420
TTGTACCGTG TGAACAACAG CAATTGCCCTG CTCTCTGAGC CTCCAAGTTG GAGGGCTCAG	480
CATTTCTCTA AGGGACTAGA CCTTTCAACC TGCCCTATAG CCCTGAAATC CATCTCTGCC	540
ACAGAGACAG CTCAGGAAGC AACTTTGGGT CGTTGGTTTG ATTGAGAAGA GAAGAAAGGG	600
GCAGAGACCC AAATGCCCTT TATAGTCTG AGCTTGGGAG AGGAGGAGGA GGTGGAGGAT	660
CTGGCCGTGA AGCTCACCTC TGGAGACTCT GAATCTCATC CAGAGCCTAC TGACCATGTC	720
CTTCAGGAAA AGAAGATGGC TCTACTGAGC TTGCTGTGCT CTACTCTGGT CTCAGAAGTA	780
AACATGAACA ATACATCTGA CCCCACCCCTG GCTGCCATTT TTGAAATCTG TCGTGAACCT	840
GCCCTCCTGG AGCCTGAGTT TATCCTCAAG GCATCTTTGT ATGCCAGGCA GCAGCTGAAC	900
GTCCGGAATG TGGCCAATAA CATCTTGGCC ATTGCTGCTT TCTTGCCGGC GTGTCGCCCC	960
CACCTGCGAC GATATTTCTG TGCCATTGTC CAGCTGCCTT CTGACTGGAT CCAGGTGGCT	1020
GAGCTTTACC AGAGCCTGGC TGAGGGAGAT AAGAATAAGC TGGTGCCCTT GCCCGCCTGT	1080

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102	AACCCTCGGA	AGCACCGGGC	CAAGAGACAC	CCCCGCCGGC	CACCCCGCTC	TCCAGGGATG	1200
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104	GAGCCTCCAT	TTTCTCACAG	ATGTTTTCCA	AGGTACATAG	GGTTTCTCAG	AGAAGAGCAG	1260
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106	AGAAAGTTTG	AGAAGGCCGG	TGATACAGTG	TCAGAGAAAA	AGAATCCTCC	AAGGTTTACC	1320
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108	CTGAAGAAGC	TGGTTCAGCG	ACTGCACATC	CACAAGCCTG	CCCAGCACGT	TCAAGCCCTG	1380
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110	CTGGGTTACA	GATACCCCTC	CAACCTACAG	CTCTTTTCTC	GAAGTCGCCT	TCCTGGGCCT	1440
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118	CGCCACCATG	AGCTCATTTCT	CCAGAGACTC	CAGCATGGGA	AGTCGGTGAT	CCACAGTCGG	1680
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122	AGAAATCAAG	CATTGCCCTT	TCCTTCGAAT	ATAACACTGA	TGAGGCGGAT	ACTAACTAGA	1800
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124	AATGAAAAGA	ACCGTCCCAG	GCGGAGGTTT	CTTTGCCACC	TAAGCCGTCA	GCAGCTTCGT	1860
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138	CTGAAGACTG	CCATCAAGCT	CCAGGCTCAA	GTCCAGGAGT	TTGATGAAAA	TGATGGATGG	2280
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140	TCCCTGAATA	CTTTTGGGAA	ATACCTGCTG	TCTCTGGCTG	GCCAAAGGGT	TCCTGTGGAC	2340
141							
142	AGGGTCATCC	TCCTTGGCCA	AAGCATGGAT	GATGGAATGA	TAAATGTGGC	CAAACAGCTT	2400
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144	TACTGGCAGC	GTGTGAATTC	CAAGTGCCTC	TTTGTGTTGGTA	TCCTCCTAAG	AAGGGTACAA	2460
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148	CTGAAGTTCA	TTGCAGAGCA	TGGGGCCTCC	CATCTTCTGG	AACATGTGGG	CCAAATGGAC	2580
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152	GAAGAGGACA	CTCCAAGCCC	CTTGGCTCCT	GTTTCCCAGC	AAGGATGGCG	CAGCATCCGG	2700

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160	GGGGAGGTGG	AGAACGCACA	GCTGTTTGTG	GGGATTCTGG	GCTCCCGTTA	TGGATACATT	2940
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162	CCCCCAGCT	ACAACCTTCC	TGACCATCCA	CACTTCCACT	GGGCCAGCA	GTACCCTTCA	3000
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164	GGGCGCTCTG	TGACAGAGAT	GGAGGTGATG	CAGTTCCTGA	ACCGGAACCA	ACGTCTGCAG	3060
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166	CCCTCTGCCC	AAGCTCTCAT	CTACTTCCGG	GATTCCAGCT	TCCTCAGCTC	TGTGCCAGAT	3120
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170	AAGAGCTACC	TAAGCAGACA	GAAAGGGATA	ACCTGCCGCA	GATACCCCTG	TGAGTGGGGG	3240
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172	GGTGTGGCAG	CTGGCCGGCC	CTATGTTGGC	GGGCTGGAGG	AGTTTGGGCA	GTTGGTTCTG	3300
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176	CCAGTGTCCA	TCCCAGACGA	TGACTTGGTC	CAGGCCACCT	TCCAGCAGCT	GCAGAAGCCA	3420
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178	CCGAGTCCTG	CCCGGCCACG	CCTTCTTCAG	GACACAGTGC	AACAGCTGAT	GCTGCCCCAC	3480
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190	CAGGTCCTGA	TCATCGATGG	GGCTGATAGG	TTAGTGGACC	AGAATGGGCA	GCTGATTTCA	3840
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192	GACTGGATCC	CAAAGAAGCT	TCCCCGGTGT	GTACACCTGG	TGCTGAGTGT	GTCTAGTGAT	3900
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194	GCAGGCCTAG	GGGAGACCCT	TGAGCAGAGC	CAGGGTGCCC	ACGTGCTGGC	CTTGGGGCCT	3960
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198	CTGGAGGAGT	CACCATTTAA	CAACCAGATG	CGACTGCTGC	TGGTGAAGCG	GGAATCAGGC	4080
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200	CGGCCGCTCT	ACCTGCGCTT	GGTCACCGAT	CACCTGAGGC	TCTTCACGCT	GTATGAGCAG	4140
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202	GTGTCTGAGA	GACTCCGGAC	CCTGCCTGCC	ACTGTCCCCC	TGCTGCTGCA	GCACATCCTG	4200
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204	AGCACACTGG	AGAAGGAGCA	CGGGCCTGAT	GTCTTCCCC	AGGCCTTGAC	TGCCCTAGAA	4260
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206	GTCACACGGA	GTGGTTTGAC	TGTGGACCAG	CTGCACGGAG	TGCTGAGTGT	GTGGCGGACA	4320
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210	TACCCCATGG	GCCCGTTTGC	CTGCCTCGTC	CAGAGTCTGC	GCAGTTTGCT	AGGGGAGGGC	4440
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212	CCTCTGGAGC	GCCCTGGTGC	CCGGCTGTGC	CTCCCTGATG	GGCCCCTGAG	AACAGCAGCT	4500
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214	AAACGTTGCT	ATGGGAAGAG	GCCAGGGCTA	GAGGACACGG	CACACATCCT	CATTGCAGCT	4560
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216	CAGCTCTGGA	AGACATGTGA	CGCTGATGCC	TCAGGCACCT	TCCGAAGTTG	CCCTCCTGAG	4620
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218	GCTCTGGGAG	ACCTGCCTTA	CCACCTGCTC	CAGAGCGGGA	ACCGTGGACT	TCTTTCGAAG	4680
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220	TTCCCTACCA	ACCTCCATGT	GGTGGCTGCA	CACTTGGAAT	TGGGTCTGGT	CTCTCGGCTC	4740
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222	TTGGAGGCCC	ATGCCCTCTA	TGCTTCTTCA	GTCCCCAAAG	AGGAACAAAA	GCTCCCCGAG	4800
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224	GCTGACGTTG	CAGTGTTTCG	CACCTTCCTG	AGGCAGCAGG	CTTCAATCCT	CAGCCAGTAC	4860
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226	CCCCGGCTCC	TGCCCCAGCA	GGCAGCCAAC	CAGCCCCTGG	ACTCACCTCT	TTGCCACCAA	4920
227							
228	GCCTCGCTGC	TCTCCCGGAG	ATGGCACCTC	CAACACACAC	TACGATGGCT	TAATAAACCC	4980
229							
230	CGGACCATGA	AAAATCAGCA	AAGCTCCAGC	CTGTCTCTGG	CAGTTTCCTC	ATCCCCTACT	5040
231							
232	GCTGTGGCCT	TCTCCACCAA	TGGGCAAAGA	GCAGCTGTGG	GCACTGCCAA	TGGGACAGTT	5100
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234	TACCTGTTGG	ACCTGAGAAC	TTGGCAGGAG	GAGAAGTCTG	TGGTGAGTGG	CTGTGATGGA	5160
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236	ATCTCTGCTT	GTTTGTTCCT	CTCCGATGAT	ACACTCTTTC	TTACTGCCTT	CGACGGGCTC	5220
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238	CTGGAGCTCT	GGGACCTGCA	GCATGGTTGT	CGGGTGCTGC	AGACTAAGGC	TCACCAGTAC	5280
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240	CAAATCACTG	GCTGCTGCCT	GAGCCCAGAC	TGCCGGCTGC	TAGCCACCGT	GTGCTTGGGA	5340
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244	AAGTCCCTGA	ACTGTGTTGC	CTTCCACCCA	GAGGGGCAGG	TAATAGCCAC	AGGCAGCTGG	5460
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246	GCTGGCAGCA	TCAGCTTCTT	CCAGGTGGAT	GGGCTCAAAG	TCACCAAGGA	CCTGGGGGCA	5520
247							
248	CCCGGAGCCT	CTATCCGTAC	CTTGGCCTTC	AATGTGCCTG	GGGGGGTTGT	GGCTGTGGGC	5580
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252	CCTGCCCACC	ATGGCTTTGT	TGCTGCTGCG	CTTTTCCTGC	ATGCGGGTTG	CCAGTTACTG	5700
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254	ACGGCTGGAG	AGGATGGCAA	GGTTCAGGTG	TGGTCAGGGT	CTCTGGGTCG	GCCCCGTGGG	5760
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256	CACCTGGGTT	CCCTTTCTCT	CTCTCCTGCC	CTCTCTGTGG	CACTCAGCCC	AGATGGTGAT	5820
257							
258	CGGGTGGCTG	TTGGATATCG	AGCGGATGGC	ATTAGGATCT	ACAAAAATCTC	TTCAGGTTCC	5880

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**SEQUENCE VERIFICATION REPORT**  
**PATENT APPLICATION US/08/951,733**

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